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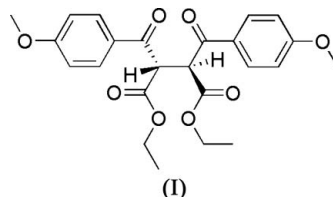
Key indicators

Single-crystal X-ray study
 $T = 292$ K
Mean $\sigma(\text{C}-\text{C}) = 0.004$ Å
 R factor = 0.056
 wR factor = 0.151
Data-to-parameter ratio = 13.7For details of how these key indicators were automatically derived from the article, see <http://journals.iucr.org/e>.

Diethyl 2,3-bis(4-methoxybenzoyl)succinate

The title molecule, $\text{C}_{24}\text{H}_{26}\text{O}_8$, has approximate twofold rotation symmetry and the crystal structure is stabilized by intermolecular $\text{C}-\text{H} \cdots \pi$ interactions.Received 28 February 2006
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Comment

1,4-Diketones are versatile intermediates for the synthesis of some natural products containing cyclopentanone and furan rings (McMurry & Meiton, 1971; Ito *et al.*, 1975, 1977). The structures of the 1,4-diketones (2*RS*,3*SR*)-diethyl 2,3-bis(3,4,5-trimethoxybenzoyl)succinate (Meng & Wu, 2005) and (2*RS*,3*SR*)-diethyl 2,3-bis(3,4-dimethoxybenzoyl)succinate (Wang *et al.*, 2005) have been reported recently and shown to be *meso* isomers. As a continuation of our interest in this area, the structure of the title compound, (I), is presented.The X-ray crystallographic analysis shows (I) to possess an approximate twofold axis (Fig. 1 and Table 1) in contrast to the above-mentioned structures. The main feature of the crystal packing are $\text{C}-\text{H} \cdots \pi$ interactions, as summarized in Table 2.

Experimental

Compound (I) was synthesized as reported previously (Wu *et al.*, 1997). Crystals appropriate for data collection were obtained by slow evaporation of a methanol–ethyl acetate (1:1 *v/v*) solution of (I).

Crystal data

 $\text{C}_{24}\text{H}_{26}\text{O}_8$
 $M_r = 442.45$
Monoclinic, $P2_1/c$
 $a = 13.079$ (5) Å
 $b = 9.611$ (4) Å
 $c = 18.572$ (7) Å
 $\beta = 102.757$ (8)°
 $V = 2276.9$ (15) Å³
 $Z = 4$ $D_x = 1.291$ Mg m⁻³
Mo $K\alpha$ radiation
Cell parameters from 3551 reflections
 $\theta = 2.4$ – 23.5°
 $\mu = 0.10$ mm⁻¹
 $T = 292$ (2) K
Block, colorless
0.30 × 0.20 × 0.18 mm

Data collection

Bruker SMART CCD area-detector diffractometer
 φ and ω scans
Absorption correction: none
11106 measured reflections
4009 independent reflections3133 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.030$
 $\theta_{\text{max}} = 25.0^\circ$
 $h = -13 \rightarrow 15$
 $k = -11 \rightarrow 9$
 $l = -16 \rightarrow 22$

Refinement

Refinement on F^2
 $R[F^2 > 2\sigma(F^2)] = 0.056$
 $wR(F^2) = 0.151$
 $S = 1.04$
 4009 reflections
 293 parameters
 H-atom parameters constrained

$$w = 1/[\sigma^2(F_o^2) + (0.0656P)^2 + 0.8894P]$$

where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.32 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.24 \text{ e } \text{\AA}^{-3}$

Table 1

Selected geometric parameters (\AA , $^\circ$).

C9–C10	1.532 (3)		
C8–C9–C10	110.75 (19)	C9–C10–C11	111.02 (18)
C8–C9–C19	108.68 (19)	C9–C10–C22	110.53 (18)
C10–C9–C19	109.62 (18)	C11–C10–C22	107.06 (18)

Table 2

Hydrogen-bond geometry (\AA , $^\circ$). $Cg1$ is the centroid of benzene ring C2–C7.

$D-H\cdots A$	$D-H$	$H\cdots A$	$D\cdots A$	$D-H\cdots A$
C14–H14 $\cdots Cg1^i$	0.93	2.92	3.743 (3)	148
C16–H16 $\cdots Cg1^{ii}$	0.93	3.00	3.689 (4)	132

Symmetry codes: (i) $x, -y + \frac{3}{2}, z - \frac{1}{2}$; (ii) $x, -y + \frac{1}{2}, z - \frac{1}{2}$.

The methyl H atoms were constrained to an ideal geometry, with C–H distances of 0.96 \AA and $U_{\text{iso}}(\text{H}) = 1.5U_{\text{eq}}(\text{C})$, but each group was allowed to rotate freely about its C–C bond. All other H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C–H distances of 0.96 \AA for aromatic and 0.97 \AA for methylene H atoms, and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C})$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT-Plus* (Bruker, 2001); data reduction: *SAINT-Plus*; program(s) used to

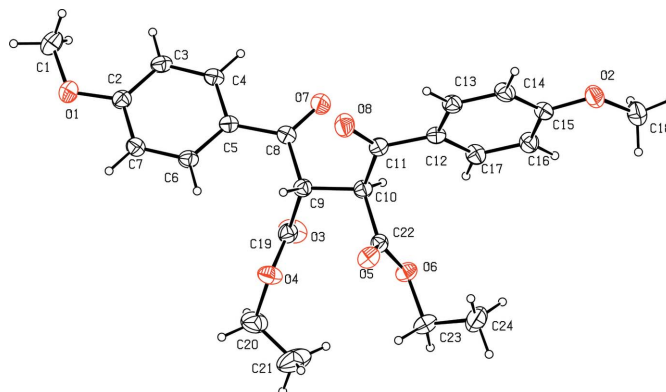


Figure 1

View of (I), showing 50% probability displacement ellipsoids and the atom-numbering scheme.

solve structure: *SHELXS97* (Sheldrick, 1997); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *PLATON* (Spek, 2003); software used to prepare material for publication: *PLATON*.

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